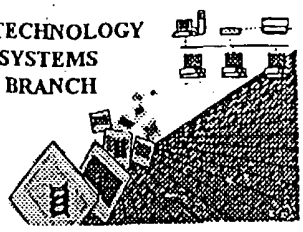
BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/648,692A
Source: IFW16 -
Date Processed by STIC: 4/22/04 -

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221
Effective 12/13/03: **TELEPHONE: 571-272-2510; FAX: 571-273-0221**

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , **EFS Submission User Manual - ePAVE)**
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. **Hand Carry directly to (EFFECTIVE 12/01/03):**
   U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. **Federal Express, United Parcel Service, or other delivery service to:** U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 09/648,692A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1	Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2	Invalid Line Length	The rules require that a line **not exceed** 72 characters in length. This includes white spaces.
3	Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.
4 ✓	Non-ASCII	The submitted file was **not** saved in ASCII(DOS) **text**, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**
5	Variable Length	Sequence(s)____contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6	PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**
7	Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.
8	Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. **If intentional**, please insert the following lines for **each** skipped sequence. <210> sequence id number <400> sequence id number 000
9	Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.
10 ✓	Invalid <213> Response	Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence
11	Use of <220>	Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
12	PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13	Misuse of n/Xaa	"n" can **only** represent a single nucleotide; "Xaa" can **only** represent a single amino acid

IFW16

RAW SEQUENCE LISTING DATE: 04/23/2004
PATENT APPLICATION: US/09/648,692A TIME: 11:49:45

Input Set : A:\Seqlist.txt
Output Set: N:\CRF4\04232004\I648692A.raw

pp 1, 3-5

Does Not Comply
Corrected Diskette Needed

see item 4 on Error Summary Sheet

```
  4 <110> APPLICANT: Dolly, James Oliver
  5       Li, Yan
  6       Chan, C.K.
  7       Aoki, Kei Roger
  9 <120> TITLE OF INVENTION: Activatable Recombinant Neurotoxins
 12 <130> FILE REFERENCE: 17311(BO)
 14 <140> CURRENT APPLICATION NUMBER: 09/648,692A
 15 <141> CURRENT FILING DATE: 2000-08-25
 17 <150> PRIOR APPLICATION NUMBER: 60/150,710
 18 <151> PRIOR FILING DATE: 1999-08-25
 20 <160> NUMBER OF SEQ ID NOS: 29
 22 <170> SOFTWARE: FastSEQ for Windows Version 4.0
```

ERRORED SEQUENCES

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 91 <210> SEQ ID NO: 7
 92 <211> LENGTH: 65
 93 <212> TYPE: PRT
 94 <213> ORGANISM: Artificial Sequence
 96 <220> FEATURE:
 97 <223> OTHER INFORMATION: Engineered Intrachain loop region for C. tetani
 98       toxin
100 <400> SEQUENCE: 7
101 Ser Lys Leu Ile Gly Leu Cys Lys Lys Ile Ile Pro Pro Thr Asn Ile
102  1               5                  10                  15
103 Arg Glu Asn Leu Tyr Asn Arg Thr Ala Gly Glu Lys Leu Tyr Asp Asp
104             20                  25                  30
105 Asp Asp Lys Asp Arg Trp Gly Ser Ser Arg Ser Leu Thr Asp Leu Gly
E--> 106
     35                  40                  45                  Gly Glu Leu Cys Ile
130 <210> SEQ ID NO: 10
131 <211> LENGTH: 4017
132 <212> TYPE: DNA
133 <213> ORGANISM: Clostridium botulinum
135 <400> SEQUENCE: 10
E--> 136 gaattcaagt agtagataat aaaaataatg ccacagattt ttattattaa taatgatata     60
137 tttatctcta actgtttaac tttaacttat aacaatgtaa atatatattt gtctataaaa     120
138 aatcaagatt acaattgggt tatatgtgat cttaatcatg atataccaaa aaagtcatat     180
139 ctatggatat taaaaaatat ataaatttaa aattaggaga tgctgtatat gccaaaaatt     240
140 aatagtttta attataatga tcctgttaat gatagaacaa ttttatatat taaaccaggc     300
141 ggttgtcaag aattttataa atcatttaat attatgaaaa atatttggat aattccagag     360
142 agaaatgtaa ttggtacaac cccccaagat tttcatccgc ctacttcatt aaaaaatgga     420
143 gatagtagtt attatgaccc taattattta caaagtgatg aagaaaagga tagattttta     480
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RAW SEQUENCE LISTING DATE: 04/23/2004
PATENT APPLICATION: US/09/648,692A TIME: 11:49:45

Input Set : A:\Seqlist.txt
Output Set: N:\CRF4\04232004\I648692A.raw

144 aaaatagtca caaaaatatt taatagaata aataataatc tttcaggagg gattttatta 540
145 gaagaactgt caaaagctaa tccatattta gggaatgata atactccaga taatcaattc 600
146 catattggtg atgcatcagc agttgagatt aaattctcaa atggtagcca agacatacta 660
147 ttacctaatg ttattataat gggagcagag cctgatttat ttgaaactaa cagttccaat 720
148 atttctctaa gaaataatta tatgccaagc aatcaccgtt ttggatcaat agctatagta 780
149 acattctcac ctgaatattc ttttagattt aatgataatt gtatgaatga atttattcaa 840
150 gatcctgctc ttacattaat gcatgaatta atacattcat tacatggact atatggggct 900
151 aaagggatta ctacaaagta tactataaca caaaaacaaa atcccctaat aacaaatata 960
152 agaggtacaa atattgaaga attcttaact tttggaggta ctgatttaaa cattattact 1020
153 agtgctcagt ccaatgatat ctatactaat cttctagctg attataaaaa aatagcgtct 1080
154 aaacttagca aagtacaagt atctaatcca ctacttaatc cttataaaga tgtttttgaa 1140
155 gcaaagtatg gattagataa agatgctagc ggaatttatt cggtaaatat aaacaaattt 1200
156 aatgatattt ttaaaaaatt atacagcttt acggaatttg atttacgaac taaatttcaa 1260
157 gttaaatgta ggcaaactta tattggacag tataaatact tcaaactttc aaacttgtta 1320
158 aatgattcta tttataatat atcagaaggc tataatataa ataatttaaa ggtaaatttt 1380
159 agaggacaga atgcaaattt aaatcctaga attattacac caattacagg tagaggacta 1440
160 gtaaaaaaaa tcattagatt ttgtaaaaat attgtttctg taaaaggcat aaggaaatca 1500
161 atatgtatcg aaataaataa tggtgagtta ttttttgtgg cttccgagaa tagttataat 1560
162 gatgataata taaatactcc taaagaaatt gacgatacag taacttcaaa taataattat 1620
163 gaaaatgatt tagatcaggt tattttaaat tttaatagtg aatcagcacc tggactttca 1680
164 gatgaaaaat taaatttaac tatccaaaat gatgcttata taccaaaata tgattctaat 1740
165 ggaacaagtg atatagaaca acatgatgtt aatgaactta atgtattttt ctatttagat 1800
166 gcacagaaag tgcccgaagg tgaaaataat gtcaatctca cctcttcaat tgatacagca 1860
167 ttattagaac aacctaaaat atatacattt ttttcatcag aatttattaa taatgtcaat 1920
168 aaacctgtgc aagcagcatt atttgtaagc tggatacaac aagtgttagt agattttact 1980
169 actgaagcta accaaaaaag tactgttgat aaaattgcag atatttctat agttgttcca 2040
170 tatataggtc ttgctttaaa tataggaaat gaagcacaaa aaggaaattt taaagatgca 2100
171 cttgaattat taggagcagg tattttatta gaatttgaac ccgagctttt aattcctaca 2160
172 attttagtat tcacgataaa atctttttta ggttcatctg ataataaaaa taaagttatt 2220
173 aaagcaataa ataatgcatt gaaagaaaga gatgaaaaat ggaaagaagt atatagtttt 2280
174 atagtatcga attggatgac taaaattaat acacaattta ataaaagaaa agaacaaatg 2340
175 tatcaagctt tacaaaatca agtaaatgca attaaaacaa taatagaatc taagtataat 2400
176 agttatactt tagaggaaaa aaatgagctt acaaataaat atgatattaa gcaaatagaa 2460
177 aatgaactta atcaaaaggt ttctatagca atgaataata tagacaggtt cttaactgaa 2520
178 agttctatat cctatttaat gaaaataata aatgaagtaa aaattaataa attaagagaa 2580
179 tatgatgaga atgtcaaaac gtatttattg aattatatta tacaacatgg atcaatcttg 2640
180 ggagagagtc agcaagaact aaattctatg gtaactgata ccctaaataa tagtattcct 2700
181 tttaagcttt cttcttatac agatgataaa attttaattt catattttaa taaattcttt 2760
182 aagagaatta aaagtagttc agttttaaat atgagatata aaaatgataa atacgtagat 2820
183 acttcaggat atgattcaaa tataaatatt aatggagatg tatataaata tccaactaat 2880
184 aaaaatcaat ttggaatata taatgataaa cttagtgaag ttaatatatc tcaaaatgat 2940
185 tacattatat atgataataa atataaaaat tttagtatta gtttttgggt aagaattcct 3000
186 aactatgata ataagatagt aaatgttaat aatgaataca ctataataaa ttgtatgaga 3060
187 gataataatt caggatggaa agtatctctt aatcataatg aaataatttg gacattcgaa 3120
188 gataatcgag gaattaatca aaaattagca tttaactatg gtaacgcaaa tggtatttct 3180
189 gattatataa ataagtggat ttttgtaact ataactaatg atagattagg agattctaaa 3240
190 ctttatatta atggaaattt aatagatcaa aaatcaattt taaatttagg taatattcat 3300
191 gttagtgaca atatattatt taaaatagtt aattgtagtt atacaagata tattggtatt 3360
192 agatatttta atatttttga taaagaatta gatgaaacag aaattcaaac tttatatagc 3420

RAW SEQUENCE LISTING DATE: 04/23/2004
PATENT APPLICATION: US/09/648,692A TIME: 11:49:45

Input Set : A:\Seqlist.txt
Output Set: N:\CRF4\04232004\I648692A.raw

193 aatgaaccta atacaaatat tttgaaggat ttttggggaa attatttgct ttatgacaaa 3480
194 gaatactatt tattaaatgt gttaaaacca aataacttta ttgataggag aaaagattct 3540
195 actttaagca ttaataatat aagaagcact attcttttag ctaatagatt atatagtgga 3600
196 ataaaagtta aaatacaaag agttaataat agtagtacta acgataatct tgttagaaag 3660
197 aatgatcagg tatatattaa ttttgtagcc agcaaaactc acttatttcc attatatgct 3720
198 gatacagcta ccacaaataa agagaaaaca ataaaaatat catcatctgg caatagattt 3780
199 aatcaagtag tagttatgaa ttcagtagga aattgtacaa tgaattttaa aaataataat 3840
E--> 200
ggaaataata ttgggttgtt aggtttcaag gcagatactg tcgttgctag tacttggtat 3900tatacacata
243 <210> SEQ ID NO: 15
244 <211> LENGTH: 5
245 <212> TYPE: PRT
246 <213> ORGANISM: Site
248 <220> FEATURE:
249 <221> NAME/KEY: SITE
250 <222> LOCATION: (1)...(5)
251 <223> OTHER INFORMATION: protease cleavage site
253 <400> SEQUENCE: 15
E--> 254 Asp Asp Asp Asp Lys 1 5
271 <210> SEQ ID NO: 17
272 <211> LENGTH: 5
273 <212> TYPE: PRT
274 <213> ORGANISM: Clostridium species
276 <220> FEATURE:
277 <221> NAME/KEY: ZN_FING
278 <222> LOCATION: (1)...(5)
279 <223> OTHER INFORMATION: Xaa=any amino acid
281 <400> SEQUENCE: 17
E--> 282 His Glu Xaa Xaa His 1 5
295 <210> SEQ ID NO: 19
296 <211> LENGTH: 22
297 <212> TYPE: PRT
298 <213> ORGANISM: Artificial Sequence
300 <220> FEATURE:
301 <223> OTHER INFORMATION: Linker
303 <400> SEQUENCE: 19
E--> 304
Met Gly Gly Ser His His His His His His Gly Met Ala Ser Met Thr 1 5
306 <210> SEQ ID NO: 20
307 <211> LENGTH: 19
308 <212> TYPE: PRT
309 <213> ORGANISM: Clostridium botulinum
311 <400> SEQUENCE: 20
E--> 312
Ser Leu Thr Asp Leu Gly Gly Glu Leu Cys Ile Lys Ile Lys Asn Glu 1 5
325 <210> SEQ ID NO: 22
326 <211> LENGTH: 7
327 <212> TYPE: PRT
328 <213> ORGANISM: Artificial Sequence
330 <220> FEATURE:
331 <221> NAME/KEY: SITE
332 <222> LOCATION: (2)...(3)

see item 4 on Error Summary Sheet
invalid <213> response
see item 10 on Error Summary Sheet (same error in sequence 16)
item 4
item 4
item 4
item 4

RAW SEQUENCE LISTING DATE: 04/23/2004
PATENT APPLICATION: US/09/648,692A TIME: 11:49:45

Input Set : A:\Seqlist.txt
Output Set: N:\CRF4\04232004\I648692A.raw

333 <223> OTHER INFORMATION: Xaa=any amino acid
335 <223> OTHER INFORMATION: Protease cleavage site
W--> 337 <400> 22
E--> 338 Glu Xaa Xaa Tyr Ser Gln Ser 1 5 item 4
340 <210> SEQ ID NO: 23
341 <211> LENGTH: 7
342 <212> TYPE: PRT
343 <213> ORGANISM: Artificial Sequence
345 <220> FEATURE:
346 <221> NAME/KEY: SITE
347 <222> LOCATION: (2)...(3)
348 <223> OTHER INFORMATION: Xaa=any amino acid
350 <221> NAME/KEY: SITE
351 <222> LOCATION: (5)...(5)
352 <223> OTHER INFORMATION: Xaa=any amino acid
354 <223> OTHER INFORMATION: Protease cleavage site
W--> 356 <400> 23
E--> 357 Glu Xaa Xaa Tyr Xaa Gln Gly 1 5 item 4
359 <210> SEQ ID NO: 24
360 <211> LENGTH: 18
361 <212> TYPE: PRT
362 <213> ORGANISM: Artificial Sequence
364 <220> FEATURE:
365 <223> OTHER INFORMATION: Translated PCR fragment item 4
367 <400> SEQUENCE: 24
E--> 368
Met Arg Gly Ser His His His His His His Gly Ser Pro Lys Ile Asn 1 5

RAW SEQUENCE LISTING DATE: 04/23/2004
PATENT APPLICATION: US/09/648,692A TIME: 11:44:32

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\04232004\I648692A.raw

181 agttctatat cctatttaat gaaaataata aatgaagtaa aaattaataa attaagagaa 2580
182 tatgatgaga atgtcaaaac gtatttattg aattatatta tacaacatgg atcaatcttg 2640
183 ggagagagtc agcaagaact aaattctatg gtaactgata ccctaaataa tagtattcct 2700
184 tttaagcttt cttcttatac agatgataaa attttaattt catattttaa taaattcttt 2760
185 aagagaatta aaagtagttc agttttaaat atgagatata aaaatgataa atacgtagat 2820
186 acttcaggat atgattcaaa tataaatatt aatggagatg tatataaata tccaactaat 2880
187 aaaaatcaat ttggaatata taatgataaa cttagtgaag ttaatatatc tcaaaatgat 2940
188 tacattatat atgataataa atataaaaat tttagtatta gtttttgggt aagaattcct 3000
189 aactatgata ataagatagt aaatgttaat aatgaataca ctataataaa ttgtatgaga 3060
190 gataataatt caggatggaa agtatctctt aatcataatg aaataatttg gacattcgaa 3120
191 gataatcgag gaattaatca aaaattagca tttaactatg gtaacgcaaa tggtatttct 3180
192 gattatataa ataagtggat ttttgtaact ataactaatg atagattagg agattctaaa 3240
193 ctttatatta atggaaattt aatagatcaa aaatcaattt taaatttagg taatattcat 3300
194 gttagtgaca atatattatt taaaatagtt aattgtagtt atacaagata tattggtatt 3360
195 agatatttta atatttttga taaagaatta gatgaaacag aaattcaaac tttatatagc 3420
196 aatgaaccta atacaaatat tttgaaggat ttttggggaa attatttgct ttatgacaaa 3480
197 gaatactatt tattaaatgt gttaaaacca aataacttta ttgataggag aaaagattct 3540
198 actttaagca ttaataatat aagaagcact attcttttag ctaatagatt atatagtgga 3600
199 ataaaagtta aaatacaaag agttaataat agtagtacta acgataatct tgttagaaag 3660
200 aatgatcagg tatatattaa ttttgtagcc agcaaaactc acttatttcc attatatgct 3720
201 gatacagcta ccacaaataa agagaaaaca ataaaaatat catcatctgg caatagattt 3780
202 aatcaagtag tagttatgaa ttcagtagga aattgtacaa tgaattttaa aaataataat 3840
203 ggaaataata ttgggttgtt aggtttcaag gcagatactg tcgttgctag tacttggtat 3900
204 tatacacata tgagagatca tacaaacagc aatggatgtt tttggaactt tatttctgaa 3960
205 gaacatggat ggcaagaaaa ataaaaatta gattaaacgg ctaaagtcat aaattcc 4017

207 <210> SEQ ID NO: 11
208 <211> LENGTH: 37 Artificial
209 <212> TYPE: DNA
C--> 210 <213> ORGANISM: Artifical Sequence
212 <220> FEATURE:
213 <223> OTHER INFORMATION: PCR primer
215 <400> SEQUENCE: 11
216 cccggatccc caaaaattaa tagttttaat tataatg 37

218 <210> SEQ ID NO: 12
219 <211> LENGTH: 36
220 <212> TYPE: DNA
221 <213> ORGANISM: PCR primer invalid <213> response
223 <400> SEQUENCE: 12
224 cccctgcagt cattttcttc gccatccatg ttcttc 36

226 <210> SEQ ID NO: 13
227 <211> LENGTH: 31
228 <212> TYPE: DNA
229 <213> ORGANISM: Artificial Sequence
231 <220> FEATURE:
232 <223> OTHER INFORMATION: PCR primer
234 <400> SEQUENCE: 13
235 cagttaatac attcattaca tggactatat g 31

237 <210> SEQ ID NO: 14
238 <211> LENGTH: 26

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 04/23/2004
PATENT APPLICATION: **US/09/648,692A** TIME: 11:49:46

Input Set : **A:\Seqlist.txt**
Output Set: **N:\CRF4\04232004\I648692A.raw**

Invalid Line Length:
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:7; Line(s) 106
Seq#:10; Line(s) 200
Seq#:19; Line(s) 304
Seq#:20; Line(s) 312
Seq#:24; Line(s) 368

VERIFICATION SUMMARY DATE: 04/23/2004
PATENT APPLICATION: **US/09/648,692A** TIME: 11:49:46

Input Set : **A:\Seqlist.txt**
Output Set: **N:\CRF4\04232004\I648692A.raw**

L:106 M:252 E: No. of Seq. differs, <211> LENGTH:Input:65 Found:48 SEQ:7
L:200 M:360 E: Sequence data overflow, line data truncated, for SEQ ID#:10
L:200 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:16
L:200 M:252 E: No. of Seq. differs, <211> LENGTH:Input:4017 Found:3840 SEQ:10
L:205 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:254 M:301 E: (44) No Sequence Data was Shown, SEQ ID:15
L:254 M:252 E: No. of Seq. differs, <211> LENGTH:Input:5 Found:0 SEQ:15
L:282 M:301 E: (44) No Sequence Data was Shown, SEQ ID:17
L:282 M:252 E: No. of Seq. differs, <211> LENGTH:Input:5 Found:0 SEQ:17
L:304 M:301 E: (44) No Sequence Data was Shown, SEQ ID:19
L:304 M:252 E: No. of Seq. differs, <211> LENGTH:Input:22 Found:0 SEQ:19
L:312 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:337 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22
L:338 M:301 E: (44) No Sequence Data was Shown, SEQ ID:22
L:338 M:252 E: No. of Seq. differs, <211> LENGTH:Input:7 Found:0 SEQ:22
L:356 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:357 M:301 E: (44) No Sequence Data was Shown, SEQ ID:23
L:357 M:252 E: No. of Seq. differs, <211> LENGTH:Input:7 Found:0 SEQ:23
L:368 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:430 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:29